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STATEMENT BY APPLICANT			PPLICANT	First Named Inventor	McGinnis, R.E.	
(Use as many sheets as necessary)				Art Unit	1631	
				Examiner Name	McGinnis, R.E.	
Sheet	6	of	10	Attorney Docket Number	2DLSM&R12/01	

		NON PATENT LITERATURE DOCUMENTS	
Examiner Initials*			
	CA	DELAHUNTY&NICKERSON, ET AL, Am J Hum Genet. 1996 Jun;58(6):1239-46. Testing the feasibility of DNA typing for human identification by PCR and an oligonucleotide ligation assay	
	СВ	KWOK ET AL, Genomics. 1996 Jan 1;31(1):123-6. Increasing the information content of STS-based genome maps: identifying polymorphisms in mapped STSs	
	СС	BUETOW ET AL, Nat Genet. 1999 Mar;21(3):323-5. Reliable identification of large numbers of candidate SNPs from public EST data	
	CD	PICOULT-NEWBERG & NICKERSON, ET AL, Genome Res. 1999 9: 167-174, Mining SNPs From EST Databases	
	CE	ROSES ET AL, Genomics. 1998 Nov 15;54(1):31-8. A 4-Mb high-density single nucleotide polymorphism-based map around human APOE.	
	CF	TOBE & NICKERSON, ET AL, Nucleic Acids Res. 1996 Oct 1;24(19):3728-32. Single-well genotyping of diallelic sequence variations by a two-color ELISA-based oligonucleotide ligat	
	CG	HACIA, FODOR, COLLINS, ET AL, Nucleic Acids Res. 1998 Nov 1;26(21):4975-82. Enhanced high density oligonucleotide array-based sequence analysis using modified nucleoside triph	
	СН	KNAPP, ET AL, Nucleic Acids Res. 1994 Oct 11;22(20):4167-75. Genetic Bit Analysis: a solid phase method for typing single nucleotide polymorphisms	
	CI	ZHAO, ET AL, Am J Hum Genet. 1998 Jul;63(1):225-40. Mapping of complex traits by single-nucleotide polymorphisms	
	CJ	DERISI & BROWN, Science Vol. 278, No. 5338, (1997), pp. 680-686 Exploring the Metabolic and Genetic Control of Gene Expression on a Genomic Scale	

Examiner	/Pahla Mhalay/	Date	
Signature	/Madio vvnaley/	Considered	10/13/2009

<sup>\*</sup>EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

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